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<110> BUMOL, Thomas Frank DOU, Shenshen GLASEBROOK, Andrew Lawrence GOULD, Kenneth Elliot HALE, John Edward HEUER, Josef Georg HUI, Kwan Yuk KHARITONENKOV, Alexei MIZRAHI, Jacques NA, Songqing NOBLITT, Timothy Wayne REIDY, Charles Arthur SONG, Ho Yeong WANG, Jian WU, Xiying ZUCKERMAN, Steven Harold

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<140> US 09/280,567

<141> 1999-03-30

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<151> 1998-12-22

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Thr	Pro	Thr 35	Tyr	Pro	Trp	Arg	Asp 40	Ala	Glu	Thr	Gly	Glu 45	Arg	Leu	Val	
Cys	Ala 50	Gln	Cys	Pro	Pro	_		Phe	Val	Gln	Arg 60	Pro	Cys	Arg	Arg	
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Phe	Trp	Asn	Tyr	Leu 85	Glu	Arg	Cys	Arg	Tyr 90	Cys	Asn	Val	Leu	Суs 95	Gly	
Glu	Arg	Glu	Glu 100	Glu	Ala	Arg	Ala	Cys 105	His	Ala	Thr	His	Asn 110	Arg	Ala	
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Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser Ser His Asp Thr Leu 185 Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala 200 Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile 215 Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu Ala Pro Glu 230 Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu Gln Leu Lys 245 250 Leu Arg Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp Gly Ala Leu 260 265 Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met Pro Gly Leu 275 280 Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 295 <210> 3 <211> 936 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (25)..(924) <400> 3 gctctccctg ctccagcaag gacc atg agg gcg ctg gag ggg cca ggc ctg 51 Met Arg Ala Leu Glu Gly Pro Gly Leu 99 Ser Leu Leu Cys Leu Val Leu Ala Leu Pro Ala Leu Leu Pro Val Pro 10 get gta ege gga gtg gea gaa aca eee ace tae eee tgg egg gae gea 147 Ala Val Arg Gly Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala 30 40 gag aca ggg gag cgg ctg gtg tgc gcc cag tgc ccc cca ggc acc ttt Glu Thr Gly Glu Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe

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		cgc Arg														291
		aac Asn														339
		acc Thr			_	_	_	_	_	_						387
		ggt Gly														435
		gcc Ala 140														483
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		cgc Arg		_	_	_	_		_	_						579
		tcc Ser														627
		agg Arg														675
		gct Ala 220														723
_	_	ctc Leu		_	_				-	_			_			771
		gcc Ala														819
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Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro Cys Arg Arg 50 55 60

Asp Ser Pro Thr Thr Cys Gly Pro Cys Pro Pro Arg His Tyr Thr Gln 65 70 75 80

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Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu 115 120 125

His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro 130 135 140

Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr Phe Ser Ala 145 150 155 160

Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala 165 170 175

Leu Gly Leu Ala Leu Ile Val Pro Gly Ser Ser Ser His Asp Thr Leu 180 185 190

Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val Pro Gly Ala 195 200 205

Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe Gln Asp Ile 210 215 220

Ser Ile Lys Arg Leu Gln Arg Leu Gln Ala Leu Glu Ala Pro Glu

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		-							_	ggc Gly	_	_		_	_		144
										cgc Arg							192
										cgg Arg							240
										ttc Phe 90							288
	tgc Cys	ttg Leu	gag Glu	cac His 100	gca Ala	tcg Ser	tgt Cys	cca Pro	cct Pro 105	ggt Gly	gcc Ala	ggc Gly	gtg Val	att Ile 110	gcc Ala	ccg Pro	336

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	_	_	_	_		_	gcc Ala										480
							tgc Cys										528
							gag Glu			_		_			_		576
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Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr 120 Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser His 155 Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val 170 Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe 185 Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu 195 200 Ala Pro Glu Gly Trp Gly Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu 215 Gln Leu Lys Leu Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp 225 230 Gly Ala Leu Leu Val Arg Leu Leu Gln Ala Leu Arg Val Ala Arg Met 245 250 Pro Gly Leu Glu Arg Ser Val Arg Glu Arg Phe Leu Pro Val His 260 265 <210> 7 <211> 825 <212> DNA <213> Homo sapiens <220> <221> CDS <222> (1)..(813) gtg gca gaa aca ccc acc tac ccc tgg cgg gac gca gag aca ggg gag Val Ala Glu Thr Pro Thr Tyr Pro Trp Arg Asp Ala Glu Thr Gly Glu

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Arg Leu Val Cys Ala Gln Cys Pro Pro Gly Thr Phe Val Gln Arg Pro

tgc cgc cga gac agc ccc acg acg tgt ggc ccg tgt cca ccg cgc cac

48

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		tgc Cys															240
		cgt Arg	_	_	_	_	_							_			288
/	_	ttg Leu			-										-	-	336
\		acc Thr															384
		tca Ser 130															432
	_	acg Thr	_	_			-										480
	_	acc Thr	_	_		_	_						_			_	528
		gga Gly															576
		gac Asp															624
		ccg Pro 210														ttg Leu	672
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Tyr Thr Gln Phe Trp Asn Tyr Leu Glu Arg Cys Arg Tyr Cys Asn Val 50 55 60

Leu Cys Gly Glu Arg Glu Glu Glu Ala Arg Ala Cys His Ala Thr His 65 70 75 80

Asn Arg Ala Cys Arg Cys Arg Thr Gly Phe Phe Ala His Ala Gly Phe
85 90 95

Cys Leu Glu His Ala Ser Cys Pro Pro Gly Ala Gly Val Ile Ala Pro 100 105 110

Gly Thr Pro Ser Gln Asn Thr Gln Cys Gln Pro Cys Pro Pro Gly Thr 115 120 125

Phe Ser Ala Ser Ser Ser Ser Glu Gln Cys Gln Pro His Arg Asn 130 135 140

Cys Thr Ala Leu Gly Leu Ala Leu Asn Val Pro Gly Ser Ser His 145 150 155 160

Asp Thr Leu Cys Thr Ser Cys Thr Gly Phe Pro Leu Ser Thr Arg Val 165 170 175

Pro Gly Ala Glu Glu Cys Glu Arg Ala Val Ile Asp Phe Val Ala Phe 180 185 190

Gln Asp Ile Ser Ile Lys Arg Leu Gln Arg Leu Leu Gln Ala Leu Glu 195 200 205

Ala Pro Glu Gly Trp Ala Pro Thr Pro Arg Ala Gly Arg Ala Ala Leu 210 215 220

Gln Leu Lys Leu Arg Arg Leu Thr Glu Leu Leu Gly Ala Gln Asp 225 230 235 240

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